

Letter

# Total Synthesis of (+)-Prunustatin A: Utility of Organotrifluoroborate-Mediated Prenylation and Shiina MNBA Esterification and Macrolactonization To Avoid a Competing Thorpe–Ingold Effect Accelerated Transesterification

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**Supporting Information** 



**ABSTRACT:** A convergent total synthesis of (+)-prunustatin A is described through the assembly of two key fragments and a macrolactonization. Shiina MNBA couplings were used for the formation of each of the four ester bonds in the tetralactone ring, including the key macrocyclization which was essential to minimize competing Thorpe–Ingold accelerated transesterification. Other key steps included an organoboron-based prenylation using potassium prenyltrifluoroborate and a carbonyldiimidazole-mediated coupling to form the salicylamide.

 $\mathbf{P}$  runustatin A (1) is a natural product isolated in 2005 from a soil sample collected on the second s in Okinawa, Japan, and determined by Shin-ya to be a potent anticancer agent against hypoxic fibrosarcoma cells.<sup>1</sup> It is produced by the bacterial species Streptomyces violaceoniger, which belongs to a family of biocontrol agents used against phytopathogenic fungi.<sup>2</sup> Structurally, 1 is a member of the antimycin depsipeptide class of compounds<sup>3</sup> that includes (+)-SW-163A  $\mathbf{2}^{4}$  JBIR-06,<sup>1e</sup> JBIR-52,<sup>5</sup> neoantimycin  $\mathbf{3}^{6}$  the parent antimycin family of compounds 4,<sup>6b,7</sup> and respirantin  $5^8$ (Figure 1). They share an aromatic 3-formamidosalicylate headgroup attached via an amide bond to an L-threonine unit embedded within a di-, tri-, or tetralactone macrocyclic ring (9-, 12-, 15-, or 18-membered). Other antimycin depsipeptides incorporate different aromatic head groups, with the entire family related through a nonribosomal peptide synthetase (NRPS)-polyketide synthase (PKS) biosynthetic pathway. Over the years, there has been significant interest in this class of depsipeptides due to their anticancer, antifungal, and immunosuppressant properties. The anticancer properties of (+)-prunustatin A are believed to stem from its ability to downregulate the expression of GRP78, which is a glucoseregulated molecular chaperone that plays an important role in the survival of tumors subjected to chemotherapy. The inhibition of GRP78 expression may be a secondary effect of blocking the mitochondrial ATP production.<sup>5</sup>

The structural motifs present in these molecules pose numerous synthetic challenges, especially the formation of the macrocyclic rings. Our previous work in this class of natural



Figure 1. Structures of (+)-prunustatin A (1) and related natural products 2-5 belonging to the antimycin class of compounds.

products has included the synthesis of antimycin  $A_{1b}$  (2014)<sup>10</sup> as well as respirantin and kitastatin (2014).<sup>11</sup> In the latter case, the presence of a lactam group permitted the use of a macrolactamization approach using HATU, whereas Shiina's

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Scheme 1. Retrosynthesis of (+)-Prunustatin A (1)



reagent (MNBA: 2-methyl-6-nitrobenzoic anhydride)<sup>12</sup> was required for the challenging macrolactonization step needed to form the 9-membered dilactone ring present in antimycin. We now report a total synthesis of (+)-prunustatin A that addresses some of the problems encountered by other groups in the total synthesis of 1.

The major challenges associated with the synthesis of (+)-prunustatin A (1) are the formation of the 15-membered tetralactone ring, the requisite seco acid precursor, and the  $\beta$ keto- $\alpha$ -gem-dimethyl ester functionality, which is also found in JBIR-06, respirantin, kitastatin, and the subsequent attachment of the 3-formamidosalicylate headgroup. Each of the existing approaches to 1 by Kawanishi,<sup>13</sup> Usuki,<sup>14</sup> and Hale<sup>15</sup> have involved linear strategies for the formation of triester precursors, followed by macrolactonization through bond a or c formation (Scheme 1). In 2014, Kawanishi achieved the first total synthesis of 1 using a macrocyclization to form bond a in 60% yield using MNBA/DMAP at 50 °C. Attempts to remove a benzyl group from the O8 position of a C9-C15 ester precursor revealed a propensity for an undesired transesterification at the C12 position with concomitant fragmentation to a butyrolactone as a result of the Thorpe-Ingold effect of the C11 gem-dimethyl group. Usuki and Hale encountered problems in their attempts to achieve macrolactonization through bond b or d formation, respectively (Scheme 1). Hale's initial approaches utilizing a bond dmacrocyclization led only to diastereomers, thus necessitating a bond a macrocyclization strategy. Hale also reported that application of Kawanishi's high-temperature conditions occurred with problems of lower yields and reproducibility and, consequently, resorted to a reoptimization of the MNBA/ DMAP conditions at room temperature and at "ultrahigh dilution" to form the desired macrolactone in 42% yield along with some dimerized products. Usuki's initial approach utilizing a bond b macrocyclization strategy was thwarted because of undesired butyrolactone formation similar to that encountered by Kawanishi. As a result, Usuki employed a strategy employed in Pettit's respirantin 5 synthesis<sup>8b</sup> involving macrocyclization of a less sterically encumbered substrate lacking the gem-dimethyl group. Thus, macrolactonization was achieved through transesterification of a  $\beta$ -ketoester leading to bond c formation in 88% yield using 20 equiv of CuSO<sub>4</sub> in toluene at reflux. This protocol requires a subsequent  $\alpha$ methylation step to yield the gem dimethyl substituted group in 62% yield.

In summary, Kawanishi employed a linear sequence of ester bond c, d, and b formation and macrocyclization of bond a, Hale employed a linear sequence of ester bond b, c, and dformation and macrocyclization of bond a, while Usuki employed a linear sequence involving ester bond a, d, and bformation, macrocyclization of bond c, and subsequent C11 gem-dimethyl group introduction. Based upon this precedent, we were particularly interested in seeing whether a fully convergent synthesis of 1 might be possible and whether the troublesome gem-dimethyl-promoted butyrolactone formation could be avoided using an approach based upon a bond bmacrocyclization of a seco acid substrate that incorporated a pre-existing C11 gem-dimethyl group. The requisite fragments would also need to be synthesized from readily available precursors using a convergent strategy and allow for the latestage incorporation of different aromatic headgroups, preferably avoiding the use of a preformed O-protected 3formamidosalicylic acid as a coupling partner.

The quasi-symmetric nature of the tetralactone ring in 1 is such that there are four possible convergent strategies for its formation. An approach involving initial C6-C12 and C14-C4 fragment synthesis through ester bond b and d formation would require subsequent ester bond *a* and *c* formation. This is undesirable due to the very hindered nature of the ester bond *c* and the possibility of competitive decarboxylation of any C9-C12  $\beta$ -keto- $\alpha$ -gem-dimethyl ester precursor fragment. Alternatively, the convergent synthetic approach involving a latestage combination of bond d and b formation would allow for the inherently more challenging ester bond *c* formation to be incorporated more straightforwardly for the formation of the "southern" fragment 9. Therefore, we chose a strategy in which the key seco acid intermediate 7 required for the lactone bond *b* formation would be formed through an intermolecular ester bond d disconnection between two similar size fragments, 8 (C2-C7) and 9 (C9-C15) which, in turn, would be accessed through ester bond a and c formation, respectively, and use a prenyl trifluoroborate salt 11 to introduce the C11 gemdimethyl group in 9 (Scheme 1).

The "northern" fragment 8 synthesis started with the formation of 15 using a modified three-step protocol of Kirschning (Scheme 2).<sup>16</sup> Acetylation of L-lactic acid to give 13 followed by an EDC-mediated *tert*-butyl protection to give 14 in 84% yield (EDC offered yield and reaction time improvement over the traditional Steglich esterification used in the original sequence). Cleavage of the acetyl group under basic

# Scheme 2. Synthesis of "Northern" Fragment 8



conditions yielded alcohol **15** in 93% yield. The synthesis of acid **16** was achieved by TBS protection of commercially available Z-Thr.<sup>17</sup> The ester coupling between **16** and **15** using MNBA, DMAP, and triethylamime afforded **17** in 90% yield. Finally, TBAF deprotection of the TBS group yielded **8** in quantitative yield.<sup>18</sup>

The synthesis of the "southern" fragment required the use of the MOM-protected  $\alpha$ -hydroxy aldehyde **12** (Scheme 3). A diazotization of D-phenylalanine in acetic acid occurs with overall retention of stereochemistry<sup>19</sup> to give the acetate **18**, which upon prolonged treatment with thionyl chloride in methanol led to **19**. MOM protection to yield **20**<sup>20</sup> followed by ester group reduction with DIBAL and Swern oxidation gave aldehyde **12** in high yield.<sup>21</sup> Given the possible epimerization

# Scheme 3. Synthesis of Aldehyde 12 and Prenylation Reaction



issues, 12 was used immediately or stored at -78 °C for up to a week with enantiomeric purity confirmed by  $[\alpha]_{D}$ determinations. The prior success that we had achieved using organotrifluoroborate salts in the synthesis of 5 encouraged us to attempt an analogous approach for the introduction of the gem-dimethyl group. Reaction of the freshly prepared aldehyde 12 with 3.0 equiv of prenyltrifluoroborate salt 11<sup>11</sup> for 3 days at room temperature using Montmorillonite K10 clay based activation afforded 10 as a 5:1 mixture of anti and syn diastereomers (10a and 10b) with a total yield of 80%. The salt 11 is a stable, white solid that can be stored for a prolonged time with minimal decomposition offering a convenient alternative to organometallic prenylations of aldehydes and ketones under mild conditions.<sup>11</sup> The reaction time and the amount of the salt 11 required could be reduced with the use of 15 mol %  $BF_3 \cdot OEt_2$  as a catalyst, giving 10 as a 4:1 mixture of diastereomers in 83% total yield.<sup>22</sup> The structure of the major diastereomer was confirmed by X-ray crystallography as having an anti relationship consistent with a Cornforth transition-state model.<sup>10</sup>

Even though both diastereomers of 10 converge onto the same product 28 during the later oxidation step, the alcohols were separated by column chromatography (Scheme 4). TBS

### Scheme 4. Synthesis of "Southern" Fragment 9



protection of the major anti diastereomer **10a** with TBS-OTf afforded **22a**, which upon dihydroxylation with OsO<sub>4</sub>, oxidative cleavage with NaIO<sub>4</sub>, and Pinnick oxidation yielded carboxylic acid **24a** (94% over three steps). The benzyl ester protected coupling partner **25** was also synthesized using a diazotization approach from L-isoleucine in two steps. The subsequent ester coupling was particularly challenging due to the steric hindrance of the carboxylic acid **24a**, and the use of Shiina conditions using 3.0 equiv of the secondary alcohol coupling partner **25** was necessary to afford ester **26a** in good yields. After TBAF deprotection of the TBS group, oxidation with DMP gave ketone **28**. An analogous sequence using the

syn diastereomer **10b** also gave **28** in similar yields. Removal of the benzyl ester protecting group from **28** using  $H_2$  and Pd/C occurred in quantitative yield to give acid **9**.

The coupling of the two fragments was once again achieved using Shiina MNBA coupling conditions to give **29** in quantitative yield (Scheme 5). The choice of acid-labile

Scheme 5. Fragment Coupling and Double Deprotection/ Macrocyclization



protecting groups in compound 29 allowed for the sequential removal of the MOM and t-Bu ester groups using TFA in a "one-pot" fashion.<sup>23</sup> After thorough drying of the reaction mixture, the seco acid 7 was obtained in 82-90% purity (NMR). The remainder of the mixture was the butyrolactone 30 and associated degraded seco acid. This degradation is facilitated by the Thorpe-Ingold effect as noted in related, more problematic, reactions in the Kawanishi<sup>13</sup> and Usuki syntheses of 1,<sup>14</sup> as well as in Pettit's respirantin synthesis in which a pyrrolidinone equivalent of the lactone was formed.<sup>8b</sup> Seco acid 7, obtained from 29 with TFA, was subjected to macrocyclization under slow-addition, high-dilution conditions (1.2 mM) with MNBA/DMAP to afford the macrocycle 6 in 59% yield over two steps. To minimize the formation of 30 during the double deprotection, TFA was removed using cold CHCl<sub>3</sub> (11-14 °C), and the reaction mixture was dried and used immediately without exposure to temperature >20  $^\circ C$  to prevent further degradation to 30. This and proper handling of the reagents allows for macrocyclization through disconnection *b* for the first time with improved efficiency and in good yield. In comparison previous macrocycle formations were performed over ester bond a (Hale: 42% yield, Kawanishi: 60% yield) and ester bond c (Usuki: 88% yield) followed by alkylation (62% yield for  $\alpha$ -methylation step; 55% yield over two steps). The key amine macrocycle 31 was then obtained in quantitative yield after Pd/C deprotection.

The 3-formamidosalicylate headgroup is typically coupled to the amine group of the macrocycle in syntheses of 1 and other antimycin analogues using a 3-formamidosalicyic acid protected at the phenolic OH. The synthesis of these precursors usually requires a few steps including a particularly harsh N-formylation reaction with sometimes difficult purification,<sup>10</sup> followed by benzyl protection of the phenolic oxygen, amide coupling with the macrocycle using HATU or EDCI, and a deprotection step.<sup>6-8,13-15</sup> In order to avoid these problems, we used an indirect approach in which unprotected 3-nitrosalicylic acid **32** was first activated with CDI (1,1'-carbonyldiimidazole) under microwave conditions for 6 min at 85 °C, cooled to room temperature, and then coupled with the macrocycle **31** in 40 min to afford **33** in 99% yield using a simple silica plug purification (Scheme 6).





Reduction of the nitro group yielded aniline 32 in 86% yield and immediate, mild formylation using *N*-formylsaccharin 35  $(Cossy's \text{ protocol})^{24}$  yielded (+)-prunustatin A (1) in 70% yield. This efficient three-step protocol potentially allows for late-stage functionalization of either the phenol or aniline functionalities on the headgroup.

In conclusion, consideration of the quasi-symmetric nature of the tetralactone ring of (+)-prunustatin A, one of the most complex cyclic polylactones known, allowed for a convergent approach to its total synthesis using Shiina MNBA coupling conditions for each of the ester bonds, a CDI-activated amide bond formation, and an N-formylsaccharin-mediated formylation. The requisite  $\alpha$ -hydroxy acid fragments were synthesized through  $\alpha$ -amino acid diazotization chemistry, and the C11 gem-dimethyl group of the  $\beta$ -keto- $\alpha$ -gem-dimethyl ester motif installed using an organoboron-based approach using a stable prenyl trifluoroborate salt. An efficient double-deprotection/ macrolactonization allowed for ring closure to occur at an ester bond that had previously been reported to be unsuccessful.<sup>14</sup> Further studies on this class of molecules and analogues will be reported in due course.

# ASSOCIATED CONTENT

### **Supporting Information**

The Supporting Information is available free of charge on the ACS Publications website at DOI: 10.1021/acs.or-glett.8b02396.

Full experimental details and characterization data for all compounds including <sup>1</sup>H and <sup>13</sup>C NMR spectral data (PDF)

# **Accession Codes**

CCDC 1858839 contains the supplementary crystallographic data for this paper. These data can be obtained free of charge via www.ccdc.cam.ac.uk/data\_request/cif, or by emailing data request@ccdc.cam.ac.uk, or by contacting The Cam-

bridge Crystallographic Data Centre, 12 Union Road, Cambridge CB2 1EZ, UK; fax: +44 1223 336033.

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